

SEQUENCE LISTING

<110> KEITH, TIM
 LITTLE, RANDALL
 VAN EERDEWEGH, PAUL
 DUPUIS, JOSEE
 DEL MASTRO, RICHARD
 SIMON, JASON
 ALLEN, KRISTINA
 PANDIT, SUNIL

<120> ANTIBODIES AND COMPOSITIONS TO A RESPIRATORY DISEASE AND OBESITY
 RELATED PROTEIN

<130> 2976-4039US3

<140> 10/670,184
 <141> 2003-09-24

<150> 09/548,797
 <151> 2000-04-13

<150> 60/129,391
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<160> 193

<170> PatentIn version 3.3

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 <213> Homo sapiens

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cta cta ctg ctg ctg ctc tgg cca gtg cca ggc gcc ggg gtg ctt caa 96
 Leu Leu Leu Leu Leu Leu Trp Pro Val Pro Gly Ala Gly Val Leu Gln
 20 25 30

gga cat atc cct ggg cag cca gtc acc ccg cac tgg gtc ctg gat gga 144
 Gly His Ile Pro Gly Gln Pro Val Thr Pro His Trp Val Leu Asp Gly
 35 40 45

caa ccc tgg cgc acc gtc agc ctg gag gag ccg gtc tcg aag cca gac 192
 Gln Pro Trp Arg Thr Val Ser Leu Glu Glu Pro Val Ser Lys Pro Asp
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atg ggg ctg gtg gcc ctg gag gct gaa ggc cag gag ctc ctg ctt gag 240
 Met Gly Leu Val Ala Leu Glu Ala Glu Gly Gln Glu Leu Leu Leu Glu
 65 70 75 80

ctg gag aag aac cac agg ctg ctg gcc cca gga tac ata gaa acc cac 288
 Leu Glu Lys Asn His Arg Leu Leu Ala Pro Gly Tyr Ile Glu Thr His
 85 90 95

tac ggc cca gat ggg cag cca gtg gtg ctg gcc ccc aac cac acg gat 336
 Tyr Gly Pro Asp Gly Gln Pro Val Val Leu Ala Pro Asn His Thr Asp

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cat tgc cac tac caa ggg cga gta agg ggc ttc ccc gac tcc tgg gta	384																			
His Cys His Tyr Gln Gly Arg Val Arg Gly Phe Pro Asp Ser Trp Val																				
115	120																			
gtc ctc tgc acc tgc tct ggg atg agt ggc ctg atc acc ctc agc agg	432																			
Val Leu Cys Thr Cys Ser Gly Met Ser Gly Leu Ile Thr Leu Ser Arg																				
130	135																			
aat gcc agc tat tat ctg cgt ccc tgg cca ccc cgg ggc tcc aag gac	480																			
Asn Ala Ser Tyr Tyr Leu Arg Pro Trp Pro Pro Arg Gly Ser Lys Asp																				
145	150																			
ttc tca acc cac gag atc ttt cgg atg gag cag ctg ctc acc tgg aaa	528																			
Phe Ser Thr His Glu Ile Phe Arg Met Glu Gln Leu Leu Thr Trp Lys																				
165	170																			
gga acc tgt ggc cac agg gat cct ggg aac aaa cgc ggc atg acc agc	576																			
Gly Thr Cys Gly His Arg Asp Pro Gly Asn Lys Ala Gly Met Thr Ser																				
180	185																			
ctt cct ggt ggt ccc cag agc agg ggc agg cga gaa cgc cgc agg acc	624																			
Leu Pro Gly Gly Pro Gln Ser Arg Gly Arg Arg Glu Ala Arg Arg Thr																				
195	200																			
cgg aag tac ctg gaa ctg tac att gtg gca gac cac acc ctg ttc ttg	672																			
Arg Lys Tyr Leu Glu Leu Tyr Ile Val Ala Asp His Thr Leu Phe Leu																				
210	215																			
act cgg cac cga aac ttg aac cac acc aaa cag cgt ctc ctg gaa gtc	720																			
Thr Arg His Arg Asn Leu Asn His Thr Lys Gln Arg Leu Leu Glu Val																				
225	230																			
gcc aac tac gtg gac cag ctt ctc agg act ctg gac att cag gtg cgc	768																			
Ala Asn Tyr Val Asp Gln Leu Leu Arg Thr Leu Asp Ile Gln Val Ala																				
245	250																			
ctg acc ggc ctg gag gtg tgg acc gag cgg gac cgc agc cgc gtc acg	816																			
Leu Thr Gly Leu Glu Val Trp Thr Glu Arg Asp Arg Ser Arg Val Thr																				
260	265																			
cag gac gcc aac gcc acg ctc tgg gcc ttc ctg cag tgg cgc cgg ggg	864																			
Gln Asp Ala Asn Ala Thr Leu Trp Ala Phe Leu Gln Trp Arg Arg Gly																				
275	280																			
ctg tgg cgc cag cgg ccc cac gac tcc cgc cag ctg ctc acg ggc cgc	912																			
Leu Trp Ala Gln Arg Pro His Asp Ser Ala Gln Leu Leu Thr Gly Arg																				
290	295																			
gcc ttc cag ggc gcc aca gtg ggc ctg cgc ccc gtc gag ggc atg tgc	960																			
Ala Phe Gln Gly Ala Thr Val Gly Leu Ala Pro Val Glu Gly Met Cys																				
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cgc gcc gag agc tgc gga ggc gtg agc acg gac cac tgc gag ctc ccc	1008																			
Arg Ala Glu Ser Ser Gly Gly Val Ser Thr Asp His Ser Glu Leu Pro																				
325	330																			
atc ggc gcc gca gcc acc atg gcc cat gag atc ggc cac agc ctc ggc	1056																			
Ile Gly Ala Ala Thr Met Ala His Glu Ile Gly His Ser Leu Gly																				
340	345																			
ctc agc cac gac ccc gac ggc tgc tgc gtg gag gct cgc gcc gag tcc	1104																			
Leu Ser His Asp Pro Asp Gly Cys Cys Val Glu Ala Ala Glu Ser																				
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	365																			

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Phe Ser Ala Cys Ser Arg Arg Gln Leu Arg Ala Phe Phe Arg Lys Gly	
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ggc ggc gct tgc ctc tcc aat gcc ccg gac ccc gga ctc ccg gtg ccg	1248
Gly Gly Ala Cys Leu Ser Asn Ala Pro Asp Pro Gly Leu Pro Val Pro	
405 410 415	
ccg gcg ctc tgc ggg aac gcc ttc gtg gaa gcg ggc gag gag tgt gac	1296
Pro Ala Leu Cys Gly Asn Gly Phe Val Glu Ala Gly Glu Glu Cys Asp	
420 425 430	
tgc ggc cct ggc cag gag tgc cgc gac ctc tgc tgc ttt gct cac aac	1344
Cys Gly Pro Gly Gln Glu Cys Arg Asp Leu Cys Cys Phe Ala His Asn	
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tgc tcg ctg cgc ccg ggg gcc cag tgc gcc cac ggg gac tgc tgc gtg	1392
Cys Ser Leu Arg Pro Gly Ala Gln Cys Ala His Gly Asp Cys Cys Val	
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Arg Cys Leu Leu Lys Pro Ala Gly Ala Leu Cys Arg Gln Ala Met Gly	
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Asp Cys Asp Leu Pro Glu Phe Cys Thr Gly Thr Ser Ser His Cys Pro	
485 490 495	
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Pro Asp Val Tyr Leu Leu Asp Gly Ser Pro Cys Ala Arg Gly Ser Gly	
500 505 510	
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Tyr Cys Trp Asp Gly Ala Cys Pro Thr Leu Glu Gln Gln Cys Gln Gln	
515 520 525	
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Leu Trp Gly Pro Asp Gly Gln Glu Val Thr Cys Arg Gly Ala Leu Ala	
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ctc ccc agt gcc cag ctg gac ctg ctt ggc ctg ggc ctg gta gag cca	1680
Leu Pro Ser Ala Gln Leu Asp Leu Leu Gly Leu Gly Leu Val Glu Pro	
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ggc acc cag tgt gga cct aga atg gtg tgc cag agc agg cgc tgc agg	1728
Gly Thr Gln Cys Gly Pro Arg Met Val Cys Gln Ser Arg Arg Cys Arg	
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His Gly Val Cys Asn Ser Asn His Asn Cys His Cys Ala Pro Gly Trp	
595 600 605	
gct cca ccc ttc tgt gac aag cca ggc ttt ggt ggc agc atg gac agt	1872
Ala Pro Pro Phe Cys Asp Lys Pro Gly Phe Gly Gly Ser Met Asp Ser	
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Cys Tyr Arg Leu Pro Gly Ala His Leu Gln Arg Cys Ser Trp Gly Cys	660	665	670	
aga agg gac cct gcg tgc agt ggc ccc aaa gat ggc cca cac agg gac				2064
Arg Arg Asp Pro Ala Cys Ser Gly Pro Lys Asp Gly Pro His Arg Asp	675	680	685	
cac ccc ctg ggc ggc gtt cac ccc atg gag ttg ggc ccc aca gcc act				2112
His Pro Leu Gly Gly Val His Pro Met Glu Leu Gly Pro Thr Ala Thr	690	695	700	
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Gly Gln Pro Trp Pro Leu Asp Pro Glu Asn Ser His Glu Pro Ser Ser	705	710	715	720
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His Pro Glu Lys Pro Leu Pro Ala Val Ser Pro Asp Pro Gln Ala Asp	725	730	735	
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Gln Val Gln Met Pro Arg Ser Cys Leu Trp	740	745		
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gccagtgaat caccggacct ccagcacctg caggcagctt ggaagtctt tcccagatg				2381
gagcttgac ccacccactc caggaaacca gagccacatt agaagtctc gagggctgga				2441
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gaggatcacc agaggccagg aggtccacac cagcctgggc aacacagcaa gacaccgcg				3041
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cta cta ctg ctg ctg ctc tgg cca gtg cca ggc gcc ggg gtg ctt caa					96
Leu Leu Leu Leu Leu Leu Trp Pro Val Pro Gly Ala Gly Val Leu Gln	20	25	30		

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Gly His Ile Pro Gly Gln Pro Val Thr Pro His Trp Val Leu Asp Gly	
35 40 45	
caa ccc tgg cgc acc gtc agc ctg gag gag ccg gtc tcg aag cca gac	192
Gln Pro Trp Arg Thr Val Ser Leu Glu Glu Pro Val Ser Lys Pro Asp	
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atg ggg ctg gtg gcc ctg gag gct gaa ggc cag gag ctc ctg ctt gag	240
Met Gly Leu Val Ala Leu Glu Ala Glu Gly Gln Leu Leu Leu Glu	
65 70 75 80	
ctg gag aag aac cac agg ctg ctg gcc cca gga tac ata gaa acc cac	288
Leu Glu Lys Asn His Arg Leu Leu Ala Pro Gly Tyr Ile Glu Thr His	
85 90 95	
tac ggc cca gat ggg cag cca gtg gtg ctg gcc ccc aac cac acg gat	336
Tyr Gly Pro Asp Gly Gln Pro Val Leu Leu Ala Pro Asn His Thr Asp	
100 105 110	
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His Cys His Tyr Gln Gly Arg Val Arg Gly Phe Pro Asp Ser Trp Val	
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Val Leu Cys Thr Cys Ser Gly Met Ser Gly Leu Ile Thr Leu Ser Arg	
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aat gcc agc tat tat ctg cgt ccc tgg cca ccc ccg ggc tcc aag gac	480
Asn Ala Ser Tyr Tyr Leu Arg Pro Trp Pro Pro Arg Gly Ser Lys Asp	
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Gly Thr Cys Gly His Arg Asp Pro Gly Asn Lys Ala Gly Met Thr Ser	
180 185 190	
ctt cct ggt ggt ccc cag agc agg ggc agg cga gaa gcg cgc agg acc	624
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225 230 235 240	
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Ala Asn Tyr Val Asp Gln Leu Leu Arg Thr Leu Asp Ile Gln Val Ala	
245 250 255	
ctg acc ggc ctg gag gtg tgg acc gag cgg gac gcg agc cgc gtc acg	816
Leu Thr Gly Leu Glu Val Trp Thr Glu Arg Asp Arg Ser Arg Val Thr	
260 265 270	
cag gac gcc aac gcc acg ctc tgg gcc ttc ctg cag tgg cgc cgg ggg	864
Gln Asp Ala Asn Ala Thr Leu Trp Ala Phe Leu Gln Trp Arg Arg Gly	
275 280 285	
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Ala	Phe	Gln	Gly	Ala	Thr	Val	Gly	Leu	Ala	Pro	Val	Glu	Gly	Met	Cys	
305					310					315					320	
cgc	gcc	gag	agc	tgc	gga	ggc	gtg	agc	acg	gac	cac	tgc	gag	ctc	ccc	1008
Arg	Ala	Glu	Ser		325	Gly	Val	Ser	Thr	Asp	His	Ser	Glu	Leu	Pro	
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atc	ggc	gcc	gca	gcc	acc	atg	gcc	cat	gag	atc	ggc	cac	agc	ctc	ggc	1056
Ile	Gly	Ala	Ala	Ala	Thr	Met	Ala	His	Glu	Ile	Gly	His	Ser	Leu	Gly	
					340				345					350		
ctc	agc	cac	gac	ccc	gac	ggc	tgc	tgc	gtg	gag	gct	gcg	gcc	gag	tcc	1104
Leu	Ser	His	Asp	Pro	Asp	Gly	Cys	Cys	Val	Glu	Ala	Ala	Ala	Glu	Ser	
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gga	ggc	tgc	gtc	atg	gct	gcg	gcc	acc	ggg	cac	ccg	ttt	ccg	cgc	gtg	1152
Gly	Gly	Cys	Val	Met	Ala	Ala	Ala	Thr	Gly	His	Pro	Phe	Pro	Arg	Val	
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ttc	agc	gcc	tgc	agc	cgc	cgc	cag	ctg	cgc	gcc	ttc	ttc	cgc	aag	ggg	1200
Phe	Ser	Ala	Cys	Ser	Arg	Arg	Gln	Leu	Arg	Ala	Phe	Phe	Arg	Lys	Gly	
385						390					395				400	
ggc	ggc	gct	tgc	ctc	tcc	aat	gcc	ccg	gac	ccc	gga	ctc	ccg	gtg	ccg	1248
Gly	Gly	Ala	Cys	Leu	Ser	Asn	Ala	Pro	Asp	Pro	Gly	Leu	Pro	Val	Pro	
					405					410					415	
ccg	gcg	ctc	tgc	ggg	aac	ggc	ttc	gtg	gaa	gcg	ggc	gag	gag	tgt	gac	1296
Pro	Ala	Leu	Cys	Gly	Asn	Gly	Phe	Val	Glu	Ala	Gly	Glu	Glu	Cys	Asp	
									425					430		
tgc	ggc	cct	ggc	cag	gag	tgc	cgc	gac	ctc	tgc	tgc	ttt	gct	cac	aac	1344
Cys	Gly	Pro	Gly	Gln	Glu	Cys	Arg	Asp	Leu	Cys	Cys	Phe	Ala	His	Asn	
							440						445			
tgc	tgc	ctg	cgc	ccg	ggg	gcc	cag	tgc	gcc	cac	ggg	gac	tgc	tgc	gtg	1392
Cys	Ser	Leu	Arg	Pro	Gly	Ala	Gln	Cys	Ala	His	Gly	Asp	Cys	Cys	Val	
						455						460				
cgc	tgc	ctg	ctg	aag	ccg	gct	gga	gcg	ctg	tgc	cgc	cag	gcc	atg	ggt	1440
Arg	Cys	Leu	Leu	Lys	Pro	Ala	Gly	Ala	Leu	Cys	Arg	Gln	Ala	Met	Gly	
465						470					475				480	
gac	tgt	gac	ctc	cct	gag	ttt	tgc	acg	ggc	acc	tcc	tcc	cac	tgt	ccc	1488
Asp	Cys	Asp	Leu	Pro	Glu	Phe	Cys	Thr	Gly	Thr	Ser	Ser	His	Cys	Pro	
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cca	gac	gtt	tac	cta	ctg	gac	ggc	tca	ccc	tgt	gcc	agg	ggc	agt	ggc	1536
Pro	Asp	Val	Tyr	Leu	Leu	Asp	Gly	Ser	Pro	Cys	Ala	Arg	Gly	Ser	Gly	
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tac	tgc	tgg	gat	ggc	gca	tgt	ccc	acg	ctg	gag	cag	cag	tgc	cag	cag	1584
Tyr	Cys	Trp	Asp	Gly	Ala	Cys	Pro	Thr	Leu	Glu	Gln	Gln	Cys	Gln	Gln	
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ctc	tgg	ggg	cct	ggc	tcc	cac	cca	gct	ccc	gag	gcc	tgt	ttc	cag	gtg	1632
Leu	Trp	Gly	Pro	Gly	Ser	His	Pro	Ala	Pro	Glu	Ala	Cys	Phe	Gln	Val	
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gtg	aac	tct	gcg	gga	gat	gct	cat	gga	aac	tgc	ggc	cag	gac	agc	gag	1680
Val	Asn	Ser	Ala	Gly	Asp	Ala	His	Gly	Asn	Cys	Gly	Gln	Asp	Ser	Glu	
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new sequence listing.txt

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Gly His Phe Leu Pro Cys Ala Gly Arg Asp Ala Leu Cys Gly Lys Leu	
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cag tgc cag ggt gga aag ccc agc ctg ctc gca ccg cac atg gtg cca	1776
Gln Cys Gln Gly Gly Lys Pro Ser Leu Leu Ala Pro His Met Val Pro	
580 585 590	
gtg gac tct acc gtt cac cta gat ggc cag gaa gtg act tgt cgg gga	1824
Val Asp Ser Thr Val His Leu Asp Gly Gln Glu Val Thr Cys Arg Gly	
595 600 605	
gcc ttg gca ctc ccc agt gcc cag ctg gac ctg ctt ggc ctg ggc ctg	1872
Ala Leu Ala Leu Pro Ser Ala Gln Leu Asp Leu Leu Gly Leu Gly Leu	
610 615 620	
gta gag cca ggc acc cag tgt gga cct aga atg gtt tgc aat agc aac	1920
Val Glu Pro Gly Thr Gln Cys Gly Pro Arg Met Val Cys Asn Ser Asn	
625 630 635 640	
cat aac tgc cac tgt gct cca ggc tgg gct cca ccc ttc tgt gac aag	1968
His Asn Cys His Cys Ala Pro Gly Trp Ala Pro Pro Phe Cys Asp Lys	
645 650 655	
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Pro Gly Phe Gly Gly Ser Met Asp Ser Gly Pro Val Gln Ala Glu Asn	
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His Leu Gln Arg Cys Ser Trp Gly Cys Arg Arg Asp Pro Ala Cys Ser	
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Gly Pro Lys Asp Gly Pro His Arg Asp His Pro Leu Gly Gly Val His	
725 730 735	
ccc atg gag ttg ggc ccc aca gcc act gga cag ccc tgg ccc ctg gac	2256
Pro Met Glu Leu Gly Pro Thr Ala Thr Gly Gln Pro Trp Pro Leu Asp	
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cct gag aac tct cat gag ccc agc agc cac cct gag aag cct ctg cca	2304
Pro Glu Asn Ser His Glu Pro Ser Ser His Pro Glu Lys Pro Leu Pro	
755 760 765	
gca gtc tcg cct gac ccc caa gca gat caa gtc cag atg cca aga tcc	2352
Ala Val Ser Pro Asp Pro Gln Ala Asp Gln Val Gln Met Pro Arg Ser	
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Cys Leu Trp	
785	
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caggaaccaca gagccacatt agaagttcct gagggctgga gaacactgct gggcacactc	2584
tcacgtctcaa taacacatca gtcccagaag caaaggtcac acagcccctg acctccctca	2644
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new sequence listing.txt

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aaaaaaaaaa	aaaaaaaaaa	aaaaaa				3390

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 <213> Homo sapiens

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cta cta ctg ctg ctg ctc tgg cca gtg cca ggc gcc ggg gtg ctt caa	96
Leu Leu Leu Leu Leu Leu Trp Pro Val Pro Gly Ala Gly Val Leu Gln	
20 25 30	
ggt gag gac gcg ggc ggg gtc ccc ctc acc ctg tgc tct gtc ttt act	144
Gly Glu Asp Ala Gly Gly Val Pro Leu Thr Leu Cys Ser Val Phe Thr	
35 40 45	
cca gga cat atc cct ggg cag cca gtc acc ccg cac tgg gtc ctg gat	192
Pro Gly His Ile Pro Gly Gln Pro Val Thr Pro His Trp Val Leu Asp	
50 55 60	
gga caa ccc tgg cgc acc gtc agc ctg gag gag ccg gtc tcg aag cca	240
Gly Gln Pro Trp Arg Thr Val Ser Leu Glu Glu Pro Val Ser Lys Pro	
65 70 75 80	
gac atg ggg ctg gtg gcc ctg gag gct gaa ggc cag gag ctc ctg ctt	288
Asp Met Gly Leu Val Ala Leu Glu Ala Glu Gly Gln Glu Leu Leu Leu	
85 90 95	
gag ctg gag aag aac cac agg ctg ctg gcc cca gga tac ata gaa acc	336
Glu Leu Glu Lys Asn His Arg Leu Leu Ala Pro Gly Tyr Ile Glu Thr	
100 105 110	
cac tac ggc cca gat ggg cag cca gtg gtg ctg gcc ccc aac cac acg	384
His Tyr Gly Pro Asp Gly Gln Pro Val Val Leu Ala Pro Asn His Thr	
115 120 125	
gat cat tgc cac tac caa ggg cga gta agg ggc ttc ccc gac tcc tgg	432
Asp His Cys His Tyr Gln Gly Arg Val Arg Gly Phe Pro Asp Ser Trp	
130 135 140	
gta gtc ctc tgc acc tgc tct ggg atg agt ggc ctg atc acc ctc agc	480
Val Val Leu Cys Thr Cys Ser Gly Met Ser Gly Leu Ile Thr Leu Ser	
145 150 155 160	
agg aat gcc agc tat tat ctg cgt ccc tgg cca ccc ccg ggc tcc aag	528
Arg Asn Ala Ser Tyr Tyr Leu Arg Pro Trp Pro Pro Arg Gly Ser Lys	

										new sequence listing.txt										
165										170										175
gac	ttc	tca	acc	cac	gag	atc	ttt	cgg	atg	gag	cag	ctg	ctc	acc	tgg	576				
Asp	Phe	Ser	Thr	His	Glu	Ile	Phe	Arg	Met	Glu	Gln	Leu	Leu	Thr	Trp					
180										185										190
aaa	gga	acc	tgt	ggc	cac	agg	gat	cct	ggg	aac	aaa	gcg	ggc	atg	acc	624				
Lys	Gly	Thr	Cys	Gly	His	Arg	Asp	Pro	Gly	Asn	Lys	Ala	Gly	Met	Thr					
195										200										205
agc	ctt	cct	ggg	ggc	ccc	cag	agc	agg	ggc	agg	cga	gaa	gcg	cgc	agg	672				
Ser	Leu	Pro	Gly	Gly	Pro	Gln	Ser	Arg	Gly	Arg	Arg	Glu	Ala	Arg	Arg					
210										215										220
acc	cgg	aag	tac	ctg	gaa	ctg	tac	att	gtg	gca	gac	acc	ctg	ttc		720				
Thr	Arg	Lys	Tyr	Leu	Glu	Leu	Tyr	Ile	Val	Ala	Asp	His	Thr	Leu	Phe					
225										230										235
ttg	act	cgg	cac	cga	aac	ttg	aac	cac	acc	aaa	cag	cgt	ctc	ctg	gaa	768				
Leu	Thr	Arg	His	Arg	Asn	Leu	Asn	His	Thr	Lys	Gln	Arg	Leu	Leu	Glu					
245										250										255
gtc	gcc	aac	tac	gtg	gac	cag	ctt	ctc	agg	act	ctg	gac	att	cag	gtg	816				
Val	Ala	Asn	Tyr	Val	Asp	Gln	Leu	Leu	Arg	Thr	Leu	Asp	Ile	Gln	Val					
260										265										270
gcg	ctg	acc	ggc	ctg	gag	gtg	tgg	acc	gag	cgg	gac	cgc	agc	cgc	gtc	864				
Ala	Leu	Thr	Gly	Leu	Glu	Val	Trp	Thr	Glu	Arg	Asp	Arg	Ser	Arg	Val					
275										280										285
acg	cag	gac	gcc	aac	gcc	acg	ctc	tgg	gcc	ttc	ctg	cag	tgg	cgc	cgg	912				
Thr	Gln	Asp	Ala	Asn	Ala	Thr	Leu	Trp	Ala	Phe	Leu	Gln	Trp	Arg	Arg					
290										295										300
ggg	ctg	tgg	gcg	cag	cgg	ccc	cac	gac	tcc	gcg	cag	ctg	ctc	acg	ggc	960				
Gly	Leu	Trp	Ala	Gln	Arg	Pro	His	Asp	Ser	Ala	Gln	Leu	Leu	Thr	Gly					
305										310										315
cgc	gcc	ttc	cag	ggc	gcc	aca	gtg	ggc	ctg	gcg	ccc	gtc	gag	ggc	atg	1008				
Arg	Ala	Phe	Gln	Gly	Ala	Thr	Val	Gly	Leu	Ala	Pro	Val	Glu	Gly	Met					
325										330										335
tgc	cgc	gcc	gag	agc	tcg	gga	ggc	gtg	agc	acg	gac	cac	tcg	gag	ctc	1056				
Cys	Arg	Ala	Glu	Ser	Ser	Gly	Gly	Val	Ser	Thr	Asp	His	Ser	Glu	Leu					
340										345										350
ccc	atc	ggc	gcc	gca	gcc	acc	atg	gcc	cat	gag	atc	ggc	cac	agc	ctc	1104				
Pro	Ile	Gly	Ala	Ala	Ala	Thr	Met	Ala	His	Glu	Ile	Gly	His	Ser	Leu					
355										360										365
ggc	ctc	agc	cac	gac	ccc	gac	ggc	tgc	tgc	gtg	gag	gct	gcg	gcc	gag	1152				
Gly	Leu	Ser	His	Asp	Pro	Asp	Gly	Cys	Cys	Val	Glu	Ala	Ala	Ala	Glu					
370										375										380
tcc	gga	ggc	tgc	gtc	atg	gct	gcg	gcc	acc	ggg	cac	ccg	ttt	ccg	cgc	1200				
Ser	Gly	Gly	Cys	Val	Met	Ala	Ala	Ala	Thr	Gly	His	Pro	Phe	Pro	Arg					
385										390										395
gtg	ttc	agc	gcc	tgc	agc	cgc	cgc	cag	ctg	cgc	gcc	ttc	ttc	cgc	aag	1248				
Val	Phe	Ser	Ala	Cys	Ser	Arg	Arg	Gln	Leu	Arg	Ala	Phe	Phe	Arg	Lys					
405										410										415
ggg	ggc	ggc	gct	tgc	ctc	tcc	aat	gcc	ccg	gac	ccc	gga	ctc	ccg	gtg	1296				
Gly	Gly	Gly	Ala	Cys	Leu	Ser	Asn	Ala	Pro	Asp	Pro	Gly	Leu	Pro	Val					
420										425										430

new sequence listing.txt

ccg	ccg	gcg	ctc	tgc	ggg	aac	ggc	ttc	gtg	gaa	gcg	ggc	gag	gag	tgt	1344
Pro	Pro	Ala	Leu	Cys	Gly	Asn	Gly	Phe	Val	Glu	Ala	Gly	Glu	Glu	Cys	
		435					440					445				
gac	tgc	ggc	cct	ggc	cag	gag	tgc	cgc	gac	ctc	tgc	tgc	ttt	gct	cac	1392
Asp	Cys	Gly	Pro	Gly	Gln	Glu	Cys	Arg	Asp	Leu	Cys	Cys	Phe	Ala	His	
	450					455					460					
aac	tgc	tcg	ctg	cgc	ccg	ggg	gcc	cag	tgc	gcc	cac	ggg	gac	tgc	tgc	1440
Asn	Cys	Ser	Leu	Arg	Pro	Gly	Ala	Gln	Cys	Ala	His	Gly	Asp	Cys	Cys	
	465				470					475				480		
gtg	cgc	tgc	ctg	ctg	aag	ccg	gct	gga	gcg	ctg	tgc	cgc	cag	gcc	atg	1488
Val	Arg	Cys	Leu	Leu	Lys	Pro	Ala	Gly	Ala	Leu	Cys	Arg	Gln	Ala	Met	
				485					490					495		
ggt	gac	tgt	gac	ctc	cct	gag	ttt	tgc	acg	ggc	acc	tcc	tcc	cac	tgt	1536
Gly	Asp	Cys	Asp	Leu	Pro	Glu	Phe	Cys	Thr	Gly	Thr	Ser	Ser	His	Cys	
			500					505					510			
ccc	cca	gac	gtt	tac	cta	ctg	gac	ggc	tca	ccc	tgt	gcc	agg	ggc	agt	1584
Pro	Pro	Asp	Val	Tyr	Leu	Leu	Asp	Gly	Ser	Pro	Cys	Ala	Arg	Gly	Ser	
		515					520					525				
ggc	tac	tgc	tgg	gat	ggc	gca	tgt	ccc	acg	ctg	gag	cag	cag	tgc	cag	1632
Gly	Tyr	Cys	Trp	Asp	Gly	Ala	Cys	Pro	Thr	Leu	Glu	Gln	Gln	Cys	Gln	
	530				535						540					
cag	ctc	tgg	ggg	cct	ggc	tcc	cac	cca	gct	ccc	gag	gcc	tgt	ttc	cag	1680
Gln	Leu	Trp	Gly	Pro	Gly	Ser	His	Pro	Ala	Pro	Glu	Ala	Cys	Phe	Gln	
	545				550					555				560		
gtg	gtg	aac	tct	gcg	gga	gat	gct	cat	gga	aac	tgc	ggc	cag	gac	agc	1728
Val	Val	Asn	Ser	Ala	Gly	Asp	Ala	His	Gly	Asn	Cys	Gly	Gln	Asp	Ser	
			565					570						575		
gag	ggc	cac	ttc	ctg	ccc	tgt	gca	ggg	agg	gat	gcc	ctg	tgt	ggg	aag	1776
Glu	Gly	His	Phe	Leu	Pro	Cys	Ala	Gly	Arg	Asp	Ala	Leu	Cys	Gly	Lys	
		580					585					590				
ctg	cag	tgc	cag	ggt	gga	aag	ccc	agc	ctg	ctc	gca	ccg	cac	atg	gtg	1824
Leu	Gln	Cys	Gln	Gly	Gly	Lys	Pro	Ser	Leu	Leu	Ala	Pro	His	Met	Val	
		595				600						605				
cca	gtg	gac	tct	acc	gtt	cac	cta	gat	ggc	cag	gaa	gtg	act	tgt	cgg	1872
Pro	Val	Asp	Ser	Thr	Val	His	Leu	Asp	Gly	Gln	Glu	Val	Thr	Cys	Arg	
		610				615					620					
gga	gcc	ttg	gca	ctc	ccc	agt	gcc	cag	ctg	gac	ctg	ctt	ggc	ctg	ggc	1920
Gly	Ala	Leu	Ala	Leu	Pro	Ser	Ala	Gln	Leu	Asp	Leu	Leu	Gly	Leu	Gly	
	625				630					635				640		
ctg	gta	gag	cca	ggc	acc	cag	tgt	gga	cct	aga	atg	gtg	tgc	cag	agc	1968
Leu	Val	Glu	Pro	Gly	Thr	Gln	Cys	Gly	Pro	Arg	Met	Val	Cys	Gln	Ser	
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agg	cgc	tgc	agg	aag	aat	gcc	ttc	cag	gag	ctt	cag	cgc	tgc	ctg	act	2016
Arg	Arg	Cys	Arg	Lys	Asn	Ala	Phe	Gln	Glu	Leu	Gln	Arg	Cys	Leu	Thr	
			660				665						670			
gcc	tgc	cac	agc	cac	ggg	gtt	tgc	aat	agc	aac	cat	aac	tgc	cac	tgt	2064
Ala	Cys	His	Ser	His	Gly	Val	Cys	Asn	Ser	Asn	His	Asn	Cys	His	Cys	
		675					680					685				
gct	cca	ggc	tgg	gct	cca	ccc	ttc	tgt	gac	aag	cca	ggc	ttt	ggt	ggc	2112
Ala	Pro	Gly	Trp	Ala	Pro	Pro	Phe	Cys	Asp	Lys	Pro	Gly	Phe	Gly	Gly	

new sequence listing.txt

690	695	700	
agc atg gac agt ggc cct gtg cag gct gaa aac cat gac acc ttc ctg			2160
Ser Met Asp Ser Gly Pro Val Gln Ala Glu Asn His Asp Thr Phe Leu			
705	710	715	720
ctg gcc atg ctc ctc agc gtc ctg ctg cct ctg ctc cca ggg gcc gcc			2208
Leu Ala Met Leu Leu Ser Val Leu Leu Pro Leu Leu Pro Gly Ala Gly			
	725	730	735
ctg gcc tgg tgt tgc tac cga ctc cca gga gcc cat ctg cag cga tgc			2256
Leu Ala Trp Cys Cys Tyr Arg Leu Pro Gly Ala His Leu Gln Arg Cys			
	740	745	750
agc tgg ggc tgc aga agg gac cct gcg tgc agt ggc ccc aaa gat gcc			2304
Ser Trp Gly Cys Arg Arg Asp Pro Ala Cys Ser Gly Pro Lys Asp Gly			
	755	760	765
cca cac agg gac cac ccc ctg ggc ggc gtt cac ccc atg gag ttg gcc			2352
Pro His Arg Asp His Pro Leu Gly Gly Val His Pro Met Glu Leu Gly			
	770	775	780
ccc aca gcc act gga cag ccc tgg ccc ctg gcc cca ggg tct cct gct			2400
Pro Thr Ala Thr Gly Gln Pro Trp Pro Leu Ala Pro Gly Ser Pro Ala			
	785	790	800
gac cat att cac aac att tac cct cca cca ttt ctc cca gac cct gag			2448
Asp His Ile His Asn Ile Tyr Pro Pro Pro Phe Leu Pro Asp Pro Glu			
	805	810	815
aac tct cat gag ccc agc agc cac cct gag aag cct ctg cca gca gtc			2496
Asn Ser His Glu Pro Ser Ser His Pro Glu Lys Pro Leu Pro Ala Val			
	820	825	830
tcg cct gac ccc caa gca gat caa gtc cag atg cca aga tcc tgc ctc			2544
Ser Pro Asp Pro Gln Ala Asp Gln Val Gln Met Pro Arg Ser Cys Leu			
	835	840	845
tgg tga gaggtagctc ctaaaatgaa cagatttaaa gacaggtggc cactgacagc			2600
Trp			
cactccagga acttgaactg caggggcaga gccagtgaat caccggacct ccagcacctg			2660
caggcagcgtt ggaagtgttc tcccgcagtg gagcttcgac ccacccactc caggaaacca			2720
gagccacact agaagttcct gagggctgga gaacactgct gggcacactc tcacgtctaa			2780
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ggttctctgag ccccaccacc caatcccagt gctacacctg aggttctgga gctcagaatc			3140
tgacagcttc tcccctatc ttgtgtgtgt ggggggacag agggaaacct ttaagaaaag			3200
ataccaaagt agaagtcmaa agaaagacat gttggctata ggcgtgtgtg ctcatgcta			3260
taatccagc actttgggaa gcyggggtag gaggatcacc agagggccags aggtccacac			3320
cagcctgggc aacacagcaa gacacgcgat ctacaaaaaa attttaaaat tagctggggc			3380
tggttggtgt taacctgagg cctagctgct caggaggctg aagcaggagg atcacttgag			3440
cctgagttca acactgcagt gagctatggt ggcaccactg cactccagcc tgggtgacag			3500
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aaaaaaaaa aaaaaaaaaa aa			3582

<210> 4
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 <212> PRT
 <213> Homo sapiens

<400> 4

new sequence listing.txt

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Met Gly Trp Arg Pro Arg Arg Ala Arg Gly Thr Pro Leu Leu Leu Leu
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Leu Leu Leu Leu Leu Leu Trp Pro Val Pro Gly Ala Gly Val Leu Gln
 20      25      30
Gly His Ile Pro Gly Gln Pro Val Thr Pro His Trp Val Leu Asp Gly
 35      40      45
Gln Pro Trp Arg Thr Val Ser Leu Glu Glu Pro Val Ser Lys Pro Asp
 50      55      60
Met Gly Leu Val Ala Leu Glu Ala Glu Gly Gln Glu Leu Leu Leu Glu
 65      70      75      80
Leu Glu Lys Asn His Arg Leu Leu Ala Pro Gly Tyr Ile Glu Thr His
 85      90      95
Tyr Gly Pro Asp Gly Gln Pro Val Val Leu Ala Pro Asn His Thr Asp
100      105      110
His Cys His Tyr Gln Gly Arg Val Arg Gly Phe Pro Asp Ser Trp Val
115      120      125
Val Leu Cys Thr Cys Ser Gly Met Ser Gly Leu Ile Thr Leu Ser Arg
130      135      140
Asn Ala Ser Tyr Tyr Leu Arg Pro Trp Pro Pro Arg Gly Ser Lys Asp
145      150      155      160
Phe Ser Thr His Glu Ile Phe Arg Met Glu Gln Leu Leu Thr Trp Lys
165      170      175
Gly Thr Cys Gly His Arg Asp Pro Gly Asn Lys Ala Gly Met Thr Ser
180      185      190
Leu Pro Gly Gly Pro Gln Ser Arg Gly Arg Arg Glu Ala Arg Arg Thr
195      200      205
Arg Lys Tyr Leu Glu Leu Tyr Ile Val Ala Asp His Thr Leu Phe Leu
210      215      220
Thr Arg His Arg Asn Leu Asn His Thr Lys Gln Arg Leu Leu Glu Val
225      230      235      240
Ala Asn Tyr Val Asp Gln Leu Leu Arg Thr Leu Asp Ile Gln Val Ala
245      250      255
Leu Thr Gly Leu Glu Val Trp Thr Glu Arg Asp Arg Ser Arg Val Thr
260      265      270
Gln Asp Ala Asn Ala Thr Leu Trp Ala Phe Leu Gln Trp Arg Arg Gly
275      280      285
Leu Trp Ala Gln Arg Pro His Asp Ser Ala Gln Leu Leu Thr Gly Arg
290      295      300
Ala Phe Gln Gly Ala Thr Val Gly Leu Ala Pro Val Glu Gly Met Cys
305      310      315      320
Arg Ala Glu Ser Ser Gly Gly Val Ser Thr Asp His Ser Glu Leu Pro
325      330      335
Ile Gly Ala Ala Ala Thr Met Ala His Glu Ile Gly His Ser Leu Gly
340      345      350

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new sequence listing.txt

Leu Ser His Asp Pro Asp Gly Cys Cys Val Glu Ala Ala Ala Glu Ser
 355 360 365
 Gly Gly Cys Val Met Ala Ala Ala Thr Gly His Pro Phe Pro Arg Val
 370 375 380
 Phe Ser Ala Cys Ser Arg Arg Gln Leu Arg Ala Phe Phe Arg Lys Gly
 385 390 395 400
 Gly Gly Ala Cys Leu Ser Asn Ala Pro Asp Pro Gly Leu Pro Val Pro
 405 410 415
 Pro Ala Leu Cys Gly Asn Gly Phe Val Glu Ala Gly Glu Glu Cys Asp
 420 425 430
 Cys Gly Pro Gly Gln Glu Cys Arg Asp Leu Cys Cys Phe Ala His Asn
 435 440 445
 Cys Ser Leu Arg Pro Gly Ala Gln Cys Ala His Gly Asp Cys Cys Val
 450 455 460
 Arg Cys Leu Leu Lys Pro Ala Gly Ala Leu Cys Arg Gln Ala Met Gly
 465 470 475 480
 Asp Cys Asp Leu Pro Glu Phe Cys Thr Gly Thr Ser Ser His Cys Pro
 485 490 495
 Pro Asp Val Tyr Leu Leu Asp Gly Ser Pro Cys Ala Arg Gly Ser Gly
 500 505 510
 Tyr Cys Trp Asp Gly Ala Cys Pro Thr Leu Glu Gln Gln Cys Gln Gln
 515 520 525
 Leu Trp Gly Pro Asp Gly Gln Glu Val Thr Cys Arg Gly Ala Leu Ala
 530 535 540
 Leu Pro Ser Ala Gln Leu Asp Leu Leu Gly Leu Gly Leu Val Glu Pro
 545 550 555 560
 Gly Thr Gln Cys Gly Pro Arg Met Val Cys Gln Ser Arg Arg Cys Arg
 565 570 575
 Lys Asn Ala Phe Gln Glu Leu Gln Arg Cys Leu Thr Ala Cys His Ser
 580 585 590
 His Gly Val Cys Asn Ser Asn His Asn Cys His Cys Ala Pro Gly Trp
 595 600 605
 Ala Pro Pro Phe Cys Asp Lys Pro Gly Phe Gly Gly Ser Met Asp Ser
 610 615 620
 Gly Pro Val Gln Ala Glu Asn His Asp Thr Phe Leu Leu Ala Met Leu
 625 630 635 640
 Leu Ser Val Leu Leu Pro Leu Leu Pro Gly Ala Gly Leu Ala Trp Cys
 645 650 655
 Cys Tyr Arg Leu Pro Gly Ala His Leu Gln Arg Cys Ser Trp Gly Cys
 660 665 670
 Arg Arg Asp Pro Ala Cys Ser Gly Pro Lys Asp Gly Pro His Arg Asp
 675 680 685
 His Pro Leu Gly Gly Val His Pro Met Glu Leu Gly Pro Thr Ala Thr
 690 695 700

new sequence listing.txt

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Gly Gln Pro Trp Pro Leu Asp Pro Glu Asn Ser His Glu Pro Ser Ser
705              710              715              720

His Pro Glu Lys Pro Leu Pro Ala Val Ser Pro Asp Pro Gln Ala Asp
              725              730              735

Gln Val Gln Met Pro Arg Ser Cys Leu Trp
              740              745

<210> 5
<211> 787
<212> PRT
<213> Homo sapiens

<400> 5

Met Gly Trp Arg Pro Arg Arg Ala Arg Gly Thr Pro Leu Leu Leu Leu
 1              5              10              15

Leu Leu Leu Leu Leu Trp Pro Val Pro Gly Ala Gly Val Leu Gln
              20              25              30

Gly His Ile Pro Gly Gln Pro Val Thr Pro His Trp Val Leu Asp Gly
              35              40              45

Gln Pro Trp Arg Thr Val Ser Leu Glu Glu Pro Val Ser Lys Pro Asp
 50              55              60

Met Gly Leu Val Ala Leu Glu Ala Glu Gly Gln Glu Leu Leu Leu Glu
 65              70              75              80

Leu Glu Lys Asn His Arg Leu Leu Ala Pro Gly Tyr Ile Glu Thr His
              85              90              95

Tyr Gly Pro Asp Gly Gln Pro Val Val Leu Ala Pro Asn His Thr Asp
100              105              110

His Cys His Tyr Gln Gly Arg Val Arg Gly Phe Pro Asp Ser Trp Val
115              120              125

Val Leu Cys Thr Cys Ser Gly Met Ser Gly Leu Ile Thr Leu Ser Arg
130              135              140

Asn Ala Ser Tyr Tyr Leu Arg Pro Trp Pro Pro Arg Gly Ser Lys Asp
145              150              155              160

Phe Ser Thr His Glu Ile Phe Arg Met Glu Gln Leu Leu Thr Trp Lys
              165              170              175

Gly Thr Cys Gly His Arg Asp Pro Gly Asn Lys Ala Gly Met Thr Ser
180              185              190

Leu Pro Gly Gly Pro Gln Ser Arg Gly Arg Arg Glu Ala Arg Arg Thr
195              200              205

Arg Lys Tyr Leu Glu Leu Tyr Ile Val Ala Asp His Thr Leu Phe Leu
210              215              220

Thr Arg His Arg Asn Leu Asn His Thr Lys Gln Arg Leu Leu Glu Val
225              230              235              240

Ala Asn Tyr Val Asp Gln Leu Leu Arg Thr Leu Asp Ile Gln Val Ala
245              250              255

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new sequence listing.txt

Leu Thr Gly	Leu Glu Val Trp	Thr	Glu Arg Asp Arg Ser Arg Val Thr
260			265 270
Gln Asp	Ala Asn Ala Thr Leu	Trp	Ala Phe Leu Gln Trp Arg Arg Gly
275		280	285
Leu Trp	Ala Gln Arg Pro His	Asp Ser	Ala Gln Leu Leu Thr Gly Arg
290		295	300
Ala Phe	Gln Gly Ala Thr Val	Gly Leu Ala	Pro Val Glu Gly Met Cys
305	310		315 320
Arg Ala	Glu Ser Ser Gly Gly	Val Ser	Thr Asp His Ser Glu Leu Pro
	325		330 335
Ile Gly	Ala Ala Ala Thr Met	Ala His	Glu Ile Gly His Ser Leu Gly
	340	345	350
Leu Ser	His Asp Pro Asp Gly	Cys Cys	Val Glu Ala Ala Glu Ser
355		360	365
Gly Gly	Cys Val Met Ala Ala	Ala Thr	Gly His Pro Phe Pro Arg Val
370		375	380
Phe Ser	Ala Cys Ser Arg Arg	Gln Leu Arg	Ala Phe Phe Arg Lys Gly
385	390		395 400
Gly Gly	Ala Cys Leu Ser Asn	Ala Pro	Asp Pro Gly Leu Pro Val Pro
	405		410 415
Pro Ala	Leu Cys Gly Asn Gly	Phe Val	Glu Ala Gly Glu Glu Cys Asp
	420	425	430
Cys Gly	Pro Gly Gln Glu Cys	Arg Asp	Leu Cys Cys Phe Ala His Asn
435		440	445
Cys Ser	Leu Arg Pro Gly Ala	Gln Cys	Ala His Gly Asp Cys Cys Val
450	455		460
Arg Cys	Leu Leu Lys Pro Ala	Gly Ala	Leu Cys Arg Gln Ala Met Gly
465	470		475 480
Asp Cys	Asp Leu Pro Glu Phe	Cys Thr	Gly Thr Ser Ser His Cys Pro
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Pro Asp	Val Tyr Leu Leu Asp	Gly Ser	Pro Cys Ala Arg Gly Ser Gly
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Tyr Cys	Trp Asp Gly Ala Cys	Pro Thr	Leu Glu Gln Gln Cys Gln Gln
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Leu Trp	Gly Pro Gly Ser His	Pro Ala	Pro Glu Ala Cys Phe Gln Val
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Val Asn	Ser Ala Gly Asp Ala	His Gly	Asn Cys Gly Gln Asp Ser Glu
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Gly His	Phe Leu Pro Cys Ala	Gly Arg	Asp Ala Leu Cys Gly Lys Leu
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Gln Cys	Gln Gly Gly Lys Pro	Ser Leu	Leu Ala Pro His Met Val Pro
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Val Asp	Ser Thr Val His Leu	Asp Gly	Gln Glu Val Thr Cys Arg Gly
595		600	605

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His Asn Cys His Cys Ala Pro Gly Trp Ala Pro Pro Phe Cys Asp Lys
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Pro Gly Phe Gly Gly Ser Met Asp Ser Gly Pro Val Gln Ala Glu Asn
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His Asp Thr Phe Leu Leu Ala Met Leu Leu Ser Val Leu Leu Pro Leu
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Leu Pro Gly Ala Gly Leu Ala Trp Cys Cys Tyr Arg Leu Pro Gly Ala
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His Leu Gln Arg Cys Ser Trp Gly Cys Arg Arg Asp Pro Ala Cys Ser
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Gly Pro Lys Asp Gly Pro His Arg Asp His Pro Leu Gly Gly Val His
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Pro Met Glu Leu Gly Pro Thr Ala Thr Gly Gln Pro Trp Pro Leu Asp
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Pro Glu Asn Ser His Glu Pro Ser Ser His Pro Glu Lys Pro Leu Pro
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Cys Leu Trp
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Gly Glu Asp Ala Gly Gly Val Pro Leu Thr Leu Cys Ser Val Phe Thr
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Pro Gly His Ile Pro Gly Gln Pro Val Thr Pro His Trp Val Leu Asp
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Gly Gln Pro Trp Arg Thr Val Ser Leu Glu Glu Pro Val Ser Lys Pro
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Asp Met Gly Leu Val Ala Leu Glu Ala Glu Gly Gln Glu Leu Leu Leu
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Glu Leu Glu Lys Asn His Arg Leu Leu Ala Pro Gly Tyr Ile Glu Thr
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His Tyr Gly Pro Asp Gly Gln Pro Val Val Leu Ala Pro Asn His Thr

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new sequence listing.txt

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22

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22

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Asp

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new sequence listing.txt

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His Gly Asp Cys Cys Val Arg Cys Leu Leu Lys Pro Ala Gly Ala Leu
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Cys Arg Gln Ala Met Gly Asp Cys Asp Leu Pro Glu Phe Cys Thr Gly
          35          40          45
Thr Ser Ser His Cys Pro Pro
 10             55

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Thr Met Ala His Glu Ile Gly His Ser Leu Gly
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gcaggagtag gctcaggaag 20

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cctctcagga gtagaggccc 20

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gcacggattc cctcctcc 18

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tcctgtgtgc ttcccata 18

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ggcctcgagt ccagtatatt                                20

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tcgcctcag cttctcag                                18

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tcacgtgggt gcctctga                                18

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ccaagcacac ttgagcgtc 19

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agccatgccc tctgcttt 18

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gagggagctc tttcccca 18

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actgcaggaa ggcccagag 19

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<400> 126

tgagggacga ccaaagaaac 20

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<400> 127

caaagtcaca caacaagcgg 20

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gaacctgagg gcaccaatta 20

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acgtgcagtg agaggtccat 20

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<400> 130

gagaggtcca tgccgaga 18

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<400> 131

aaggttcagg gtgagggttt 20

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<400> 132

ctggagcaca gtggcagtta 20

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<223> Exon 216_Q - Reverse Primer
<400> 133

tgtactggga ggtagagggc 20

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ccagaaacct gattagggg 20

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<210> 139
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<210> 145

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<211> 39
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<210> 151
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<213> Artificial

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<210> 158
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<220>
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Ser Thr Ile Ser Tyr
1 5

<210> 173
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Ser Met Val Ser Tyr
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Pro Met Val Asn Tyr
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Ser Thr Ile Asn Tyr
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14

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14


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cccacttaga taat 14

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